SEQUENCE LISTING



<110> Tew, Kenneth D. Vulevic, Bojana Chen, Zhijian

<120> Nucleic Acid Encoding Human ABCA Transporter 2 and Methods of Use Thereof

<130> FCCC.99-08US

<140> 10/088,467 <141> 2002-06-24

<150> PCT/US00/40789

<151> 2000-08-31

<150> 60/154,839

<151> 1999-09-20

<160> 36

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 8040

<212> DNA

<213> Homo sapiens

<400> 1

ccgcggcgct gaggcggcgg agcgtggccc cgccatgggc ttcctgcacc agctgcagct 60 gctgctctgg aagaacgtga cgctcaaacg ccggagcccg tgggtcctgg ccttcgagat 120 cttcatcccc ctggtgctgt tctttatcct gctggggctg cgacagaaga agcccaccat 180 240 ctccgtgaag gaagtcccct tctacacagc ggcgcccctg acgtctgccg gcatcctgcc 300 tgtcatgcaa tcgctgtgcc cggacggcca gcgagacgag ttcggcttcc tgcagtacgc 360 caactccacg gtcacgcagc tgcttgagcg cctggaccgc gtggtggagg aaggcaacct 420 gtttgaccca gcgcggccca gcctgggctc agagctcgag gccctacgcc agcatctgga ggccctcagt gcgggcccgg gcacctcggg gagccacctg gacagatcca cagtgtcttc 480 540 cttctctctg gactcggtgg ccagaaaccc gcaggagctc tggcgtttcc tgacgcaaaa cttgtcgctg cccaatagca cggcccaagc actcttggcc gcccgtgtgg acccgcccga 600 660 ggtctaccac ctgctctttg gtccctcatc tgccctggat tcacagtctg gcctccacaa gggtcaggag ccctggagcc gcctaggggg caatcccctg ttccggatgg aggagctgct 720 780 getggeteet geeeteetgg ageageteae etgeaegeeg ggeteggggg agetgggeeg 840 gatcctcact gtgcctgaga gtcagaaggg agccctgcag ggctaccggg atgctgtctg cagtgggcag gctgctgcgc gtgccaggcg cttctctggg ctgtctgctg agctccggaa 900 ccagctggac gtggccaagg tctcccagca gctgggcctg gatgccccca acggctcgga 960 ctcctcgcca caggcgccac ccccacggag gctgcaggcg cttctggggg acctgctgga 1020 tgcccagaag gttctgcagg atgtggatgt cctgtcggcc ctggccctgc tactgcccca 1080 gggtgcctgc actggccgga cccccggacc cccagccagt ggtgcgggtg gggcggccaa 1140 tggcactggg gcaggggcag tcatgggccc caacgccacc gctgaggagg gcgcaccctc 1200 tgctgcagca ctggccaccc cggacacgct gcagggccag tgctcagcct tcgtacagct 1260 1320 ctgggccggc ctgcagccca tcttgtgtgg caacaaccgc accattgaac ccgaggcgct 1380 gcggcggggc aacatgagct ccctgggctt cacgagcaag gagcagcgga acctgggcct cctcgtgcac ctcatgacca gcaaccccaa aatcctgtac gcgcctgcgg gctctgaggt 1440 cgaccgcgtc atcctcaagg ccaacgagac ttttgctttt gtgggcaacg tgactcacta 1500 tgcccaggtc tggctcaaca tctcggcgga gatccgcagc ttcctggagc agggcaggct 1560 1620 gcagcaacac ctgcgctggc tgcagcagta tgtagcagag ctgcggctgc accccgaggc 1680 actgaacctg tcactggatg agctgccgcc ggccctgaga caggacaact tctcgctgcc cagtggcatg gccctcctgc agcagctgga taccattgac aacgcggcct gcggctggat 1740

1800 ccagttcatg tccaaggtga gcgtggacat cttcaagggc ttccccgacg aggagagcat 1860 tgtcaactac accetcaace aggeetacea ggacaacgte actgtttttg ceagtgtgat 1920 cttccagacc cggaaggacg gctcgctccc gcctcacgtg cactacaaga tccgccagaa ctccagcttc accgagaaaa ccaacgagat ccgccgcgcc tactggcggc ctgggcccaa 1980 tactggcggc cgcttctact tcctctacgg cttcgtctgg atccaggaca tgatggagcg 2040 cgccatcatc gacacttttg tggggcacga cgtggtggag ccaggcagct acgtgcagat 2100 gttcccctac ccctgctaca cacgcgatga cttcctgttt gtcattgagc acatgatgcc 2160 gctgtgcatg gtgatctcct gggtctactc cgtggccatg accatccagc acatcgtggc 2220 ggagaaggag caccggctca aggaggtgat gaagaccatg ggcctgaaca acgcggtgca 2280 ctgggtggcc tggttcatca ccggctttgt gcagctgtcc atctccgtga cagcactcac 2340 2400 cgccatcctg aagtacggcc aggtgcttat gcacagccac gtggtcatca tctggctctt cctggcagtc tacgcggtgg ccaccatcat gttctgcttc ctggtgtctg tgctgtactc 2460 caaggccaag ctggcctcgg cctgcggtgg catcatctac ttcctgagct acgtgcccta 2520 catgtacgtg gcgatccgag aggaggtggc gcatgataag atcacggcct tcgagaagtg 2580 catcgcgtcc ctcatgtcca cgacggcctt tggtctgggc tctaagtact tcgcgctgta 2640 tgaggtggcc ggcgtgggca tccagtggca caccttcagc cagtccccgg tggaggggga 2700 2760 cgacttcaac ttgctcctgg ctgtcaccat gctgatggtg gacgccgtgg tctatggcat cctcacgtgg tacattgagg ctgtgcaccc aggcatgtac gggctgcccc ggccctggta 2820 cttcccactg cagaagtcct actggctggg cagtgggcgg acagaagcct gggagtggag 2880 ctggccgtgg gcacgcaccc cccgcctcag tgtcatggag gaggaccagg cctgtgccat 2940 ggagagccgg cgctttgagg agacccgtgg catggaggag gagcccaccc acctgcctct 3000 ggttgtctgc gtggacaaac tcaccaaggt ctacaaggac gacaagaagc tggccctgaa 3060 caagetgage etgaacetet aegagaacea ggtggtetee ttettgggee aeaaegggge 3120 gggcaagacc accaccatgt ccatcctgac cggcctgttc cctccaacgt cgggttccgc 3180 caccatctac gggcacgaca tccgcacgga gatggatgag atccgcaaga acctgggcat 3240 gtgcccgcag cacaatgtgc tctttgaccg gctcacggtg gaggaacacc tctggttcta 3300 ctcacggctc aagagcatgg ctcaggagga gatccgcaga gagatggaca agatgatcga 3360 ggacctggag ctctccaaca aacggcactc actggtgcag acattgtcgg gtggcatgaa 3420 3480 gegeaagetg teegtggeea tegeettegt gggeggetet egegeeatea teetggaega 3540 geccaeggeg ggegtggace cetaegegeg cegegecate tgggacetea teetgaagta 3600 caagccagge egeaceatee ttetgteeae ceaceaeatg gatgaggetg acetgettgg 3660 ggaccgcatt gccatcatct cccatgggaa gctcaagtgc tgcggctccc cgctcttcct caagggcacc tatggcgacg ggtaccgcct cacgctggtc aagcggcccg ccgagccggg 3720 gggcccccaa gagccagggc tggcatccag cccccaggt cgggccccgc tgagcagctg 3780 ctccgagctc caggtgtccc agttcatccg caagcatgtg gcctcctgcc tgctggtctc 3840 3900 agacacaagc acggagetet cetacatect geceagegag geegeeaaga agggggettt 3960 cgagegeete ttecageace tggagegeag cetggatgea etgeacetea geagettegg gctgatggac acgaccctgg aggaagtgtt cctcaaggtg tcggaggagg atcagtcgct 4020 ggagaacagt gaggccgatg tgaaggagtc caggaaggat gtgctccctg gggcggaggg 4080 cccggcgtct ggggagggtc acgctggcaa tctggcccgg tgctcggagc tgacccagtc 4140 gcaggcatcg ctgcagtcgg cgtcatctgt gggctctgcc cgtggcgacg agggagctgg 4200 ctacaccgac gtctatggcg actaccgccc cctctttgat aacccacagg acccagacaa 4260 tgtcagcctg caagaggtgg aggcagaggc cctgtcgagg gtcggccagg gcagccgcaa 4320 gctggacggc gggtggctga aggtgcgcca gttccacggg ctgctggtca aacgcttcca 4380 ctgcgcccgc cgcaactcca aggcactctt ctcccagatc ttgctgccag ccttcttcgt 4440 ctgcgtggcc atgaccgtgg ccctgtccgt cccggagatt ggtgatctgc ccccgctggt 4500 cctgtcacct tcccagtacc acaactacac ccagccccgt ggcaatttca tcccctacgc 4560 4620 caacgaggag cgccgcgagt accggctgcg gctatcgccc gacgccagcc cccagcagct 4680 cgtgagcacg ttccggctgc cgtcgggggt gggtgccacc tgcgtgctca agtctcccgc 4740 caacggctcg ctggggccca cgttgaacct gagcagcggg gagtcgcgcc tgctggcggc 4800 teggttette gacageatgt gtetggagte etteacacag gggetgeeae tgtecaattt cgtgccacc ccacctcgc ccgccccatc tgactcgcca gcgtccccgg atgaggacct 4860 4920 graggertgg aargteteer tgregerear egetgggera gaaatgtgga egteggeace 4980 ctccctgccg cgcctggtac gggagcccgt ccgctgcacc tgctctgcgc agggcaccgg 5040 cttctcctgc cccagcagtg tgggcgggca cccgccccag atgcgggtgg tcacaggcga catcctgacc gacatcaccg gccacaatgt ctctgagtac ctgctcttca cctccgaccg 5100 5160 cttccgactg caccggtatg gggccatcac ctttggaaac gtcctgaagt ccatcccagc ctcatttggc accagggccc cacccatggt gcggaagatc gcggtgcgca gggctgccca 5220 5280 ggttttctac aacaacaagg gctatcacag catgcccacc tacctcaaca gcctcaacaa cgccatcctg cgtgccaacc tgcccaagag caagggcaac ccggcggctt acggcatcac 5340 cgtcaccaac caccccatga ataagaccag cgccagcctc tccctggatt acctgctgca 5400

```
5460
gggcacggat gtcgtcatcg ccatcttcat catcgtggcc atgtccttcg tgccggccag
                                                                      5520
cttcgttgtc ttcctcgtgg ccgagaagtc caccaaggcc aagcatctgc agtttgtcag
                                                                      5580
cggctgcaac cccatcatct actggctggc gaactacgtg tgggacatgc tcaactacct
ggtccccgct acctgctgtg tcatcatcct gtttgtgttc gacctgccgg cctacacgtc
                                                                      5640
                                                                      5700
gcccaccaac ttccctgccg tcctctccct cttcctgctc tatgggtggt ccatcacgcc
catcatgtac ccggcctcct tctggttcga ggtccccagc tccgcctacg tgttcctcat
                                                                      5760
tgtcatcaat ctcttcatcg gcatcaccgc caccgtggcc accttcctgc tacagctctt
                                                                      5820
cgagcacgac aaggacctga aggttgtcaa cagttacctg aaaagctgct tcctcatttt
                                                                      5880
                                                                      5940
ccccaactac aacctgggcc acgggctcat ggagatggcc tacaacgagt acatcaacga
gtactacgcc aagattggcc agtttgacaa gatgaagtcc ccgttcgagt gggacattgt
                                                                      6000
caccegegga etggtggeea tggeggttga gggegtegtg ggetteetee tgaccateat
                                                                      6060
gtgccagtac aacttcctgc ggcggccaca gcgcatgcct gtgtctacca agcctgtgga
                                                                      6120
ggatgatgtg gacgtggcca gtgagcggca gcgagtgctc cggggagacg ccgacaatga
                                                                      6180
catggtcaag attgagaacc tgaccaaggt ctacaagtcc cggaagattg gccgtatcct
                                                                      6240
ggccgttgac cgcctgtgcc tgggtgtgcg tcctggcgag tgcttcgggc tcctgggcgt
                                                                      6300
caacggtgcg ggcaagacca gcaccttcaa gatgctgacc ggcgacgaga gcacgacggg
                                                                      6360
                                                                      6420
gggcgaggcc ttcgtcaatg gacacagcgt gctgaaggag ctgctccagg tgcagcagag
                                                                      6480
cctcggctac tgcccgcagt gtgacgcgct gttcgacgag ctcacggccc gggagcacct
gcagctgtac acgcggctgc gtgggatctc ctggaaggac gaggcccggg tggtgaagtg
                                                                      6540
ggctctggag aagctggagc tgaccaagta cgcagacaag ccggctggca cctacagcgg
                                                                      6600
cggcaacaag cggaagctct ccacggccat cgccctcatt gggtacccag ccttcatctt
                                                                      6660
cctggacgag cccaccacag gcatggaccc caaggcccgg cgcttcctct ggaacctcat
                                                                      6720
cctcgacctc atcaagacag ggcgttcagt ggtgctgaca tcacacagca tggaggagtg
                                                                      6780
cgaggcgctg tgcacgcggc tggccatcat ggtgaacggt cgcctgcggt gcctgggcag
                                                                      6840
catccagcac ctgaagaacc ggtttggaga tggctacatg atcacggtgc ggaccaagag
                                                                      6900
cagccagagt gtgaaggacg tggtgcggtt cttcaaccgc aacttcccgg aagccatgct
                                                                      6960
                                                                      7020
caaggagegg caccacacaa aggtgeagta ceageteaag teggageaea tetegetgge
                                                                      7080
ccaggtgttc agcaagatgg agcaggtgtc tggcgtgctg ggcatcgagg actactcggt
                                                                      7140
cagccagacc acactggaca atgtgttcgt gaactttgcc aagaagcaga gtgacaacct
ggagcagcag gagacggagc cgccatccgc actgcagtcc cctctcggct gcttgctcag
                                                                      7200
cctgctccgg ccccggtctg ccccacgga gctccgggca cttgtggcag acgagcccga
                                                                      7260
ggacctggac acggaggacg agggcctcat cagcttcgag gaggagcggg cccagctgtc
                                                                      7320
cttcaacacg gacacgctct gctgaccacc cagagctggg ccagggagga cacgctccac
                                                                      7380
tgaccaccca gagctgggcc agggactcaa caatggggac agaagtcccc cagtgcctgc
                                                                      7440
cagggcctgg agtggaggtt caggaccaag gggcttctgg tcctccagcc cctgtactcg
                                                                      7500
gccatgccct gcggtcactg cggttgccgc ccctaattgt gccaaaggct gacccggccc
                                                                      7560
gggctgcgta caccettgce etgetttgce ttaaageete ggggtetgee eggeeeeteg
                                                                      7620
cccctgcctg gcactgctca ccgcccaagg cgacgccggc tggaccaggc actgctggcc
                                                                      7680
tttctcctgc ccggcctcgg aaccagcttt tctctcttac gatgaaggct gatgccgaga
                                                                      7740
gcgggctgtg ggcggagctg ggtcagtccc gtatttattt tgctttgaga agaggctcct
                                                                      7800
ctggccctgc tctcctgcag ggaggtggct gtcccgcggg aagccatcag cttgggccag
                                                                      7860
                                                                      7920
ctggcaggtg gcaggaatgg agaagctgac cctgctggcc aggcaagggg ccagaccccc
                                                                      7980
cccaacccc agetgecate geteteceae ceagettgge eccetgeceg eccaectece
                                                                      8040
tgggagccgg gcctgtacat agcgcacaga tgtttgtttt aaataaataa acaaaatgtc
```

<210> 2 <211> 2436 <212> PRT <213> Homo sapiens

 Act of the control o

Asp Glu Phe Gly Phe Leu Gln Tyr Ala Asn Ser Thr Val Thr Gln Leu Leu Glu Arg Leu Asp Arg Val Val Glu Glu Gly Asn Leu Phe Asp Pro Ala Arg Pro Ser Leu Gly Ser Glu Leu Glu Ala Leu Arg Gln His Leu Glu Ala Leu Ser Ala Gly Pro Gly Thr Ser Gly Ser His Leu Asp Arg Ser Thr Val Ser Ser Phe Ser Leu Asp Ser Val Ala Arg Asn Pro Gln Glu Leu Trp Arg Phe Leu Thr Gln Asn Leu Ser Leu Pro Asn Ser Thr Ala Gln Ala Leu Leu Ala Ala Arg Val Asp Pro Pro Glu Val Tyr His Leu Leu Phe Gly Pro Ser Ser Ala Leu Asp Ser Gln Ser Gly Leu His Lys Gly Gln Glu Pro Trp Ser Arg Leu Gly Gly Asn Pro Leu Phe Arg Met Glu Glu Leu Leu Ala Pro Ala Leu Leu Glu Gln Leu Thr Cys Thr Pro Gly Ser Gly Glu Leu Gly Arg Ile Leu Thr Val Pro Glu Ser Gln Lys Gly Ala Leu Gln Gly Tyr Arg Asp Ala Val Cys Ser Gly Gln Ala Ala Ala Arg Ala Arg Arg Phe Ser Gly Leu Ser Ala Glu Leu Arg Asn Gln Leu Asp Val Ala Lys Val Ser Gln Gln Leu Gly Leu Asp Ala Pro Asn Gly Ser Asp Ser Ser Pro Gln Ala Pro Pro Pro Arg Arg Leu Gln Ala Leu Leu Gly Asp Leu Leu Asp Ala Gln Lys Val Leu Gln Asp Val Asp Val Leu Ser Ala Leu Ala Leu Leu Leu Pro Gln Gly Ala Cys Thr Gly Arg Thr Pro Gly Pro Pro Ala Ser Gly Ala Gly Gly Ala Ala Asn Gly Thr Gly Ala Gly Ala Val Met Gly Pro Asn Ala Thr Ala Glu Glu Gly Ala Pro Ser Ala Ala Ala Leu Ala Thr Pro Asp Thr Leu Gln Gly Gln Cys Ser Ala Phe Val Gln Leu Trp Ala Gly Leu Gln Pro Ile Leu Cys Gly Asn Asn Arg Thr Ile Glu Pro Glu Ala Leu Arg Arg Gly Asn Met Ser Ser Leu Gly Phe Thr Ser Lys Glu Gln Arg Asn Leu Gly Leu Leu Val His Leu Met Thr Ser Asn Pro Lys Ile Leu Tyr Ala Pro Ala Gly Ser Glu Val Asp Arg Val Ile Leu Lys Ala Asn Glu Thr Phe Ala Phe Val Gly Asn Val Thr His Tyr Ala Gln Val Trp Leu Asn Ile Ser Ala Glu Ile Arg Ser Phe Leu Glu Gln Gly Arg Leu Gln Gln His Leu Arg Trp Leu Gln Gln Tyr Val Ala Glu Leu Arg Leu His Pro Glu Ala Leu Asn Leu Ser Leu Asp Glu Leu Pro Pro Ala Leu Arg Gln Asp Asn Phe Ser Leu Pro Ser Gly Met Ala Leu Leu Gln Gln Leu Asp Thr

```
Ile Asp Asn Ala Ala Cys Gly Trp Ile Gln Phe Met Ser Lys Val Ser
                                570
               565
Val Asp Ile Phe Lys Gly Phe Pro Asp Glu Glu Ser Ile Val Asn Tyr
                               585
Thr Leu Asn Gln Ala Tyr Gln Asp Asn Val Thr Val Phe Ala Ser Val
                           600
                                               605
Ile Phe Gln Thr Arg Lys Asp Gly Ser Leu Pro Pro His Val His Tyr
                       615
                                           620
Lys Ile Arg Gln Asn Ser Ser Phe Thr Glu Lys Thr Asn Glu Ile Arg
                   630
                                       635
625
Arg Ala Tyr Trp Arg Pro Gly Pro Asn Thr Gly Gly Arg Phe Tyr Phe
                                   650
               645
Leu Tyr Gly Phe Val Trp Ile Gln Asp Met Met Glu Arg Ala Ile Ile
                                                   670
           660
                               665
Asp Thr Phe Val Gly His Asp Val Val Glu Pro Gly Ser Tyr Val Gln
                                               685
                           680
       675
Met Phe Pro Tyr Pro Cys Tyr Thr Arg Asp Asp Phe Leu Phe Val Ile
                                           700
                       695
Glu His Met Met Pro Leu Cys Met Val Ile Ser Trp Val Tyr Ser Val
                                       715
                   710
Ala Met Thr Ile Gln His Ile Val Ala Glu Lys Glu His Arg Leu Lys
                                   730
               725
Glu Val Met Lys Thr Met Gly Leu Asn Asn Ala Val His Trp Val Ala
                               745
           740
Trp Phe Ile Thr Gly Phe Val Gln Leu Ser Ile Ser Val Thr Ala Leu
        755
                            760
Thr Ala Ile Leu Lys Tyr Gly Gln Val Leu Met His Ser His Val Val
                        775
                                            780
Ile Ile Trp Leu Phe Leu Ala Val Tyr Ala Val Ala Thr Ile Met Phe
                                        795
                   790
Cys Phe Leu Val Ser Val Leu Tyr Ser Lys Ala Lys Leu Ala Ser Ala
                                    810
                805
Cys Gly Gly Ile Ile Tyr Phe Leu Ser Tyr Val Pro Tyr Met Tyr Val
                                825
            820
Ala Ile Arg Glu Glu Val Ala His Asp Lys Ile Thr Ala Phe Glu Lys
                                                845
        835
                            840
Cys Ile Ala Ser Leu Met Ser Thr Thr Ala Phe Gly Leu Gly Ser Lys
                                            860
                        855
Tyr Phe Ala Leu Tyr Glu Val Ala Gly Val Gly Ile Gln Trp His Thr
                                        875
                   870
Phe Ser Gln Ser Pro Val Glu Gly Asp Asp Phe Asn Leu Leu Ala
                                    890
               885
Val Thr Met Leu Met Val Asp Ala Val Val Tyr Gly Ile Leu Thr Trp
                                905
Tyr Ile Glu Ala Val His Pro Gly Met Tyr Gly Leu Pro Arg Pro Trp
                                               925
                            920
        915
Tyr Phe Pro Leu Gln Lys Ser Tyr Trp Leu Gly Ser Gly Arg Thr Glu
                       935
                                            940
Ala Trp Glu Trp Ser Trp Pro Trp Ala Arg Thr Pro Arg Leu Ser Val
                                        955
                    950
Met Glu Glu Asp Gln Ala Cys Ala Met Glu Ser Arg Arg Phe Glu Glu
                                    970
               965
Thr Arg Gly Met Glu Glu Pro Thr His Leu Pro Leu Val Val Cys
                               985
                                                    990
            980
Val Asp Lys Leu Thr Lys Val Tyr Lys Asp Asp Lys Lys Leu Ala Leu
                            1000
                                                1005
        995
Asn Lys Leu Ser Leu Asn Leu Tyr Glu Asn Gln Val Val Ser Phe Leu
                                            1020
                       1015
Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu Thr Gly
                   1030
                                        1035
Leu Phe Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His Asp Ile
```

	1045	1050	1055
106	0	rg Lys Asn Leu Gly 1065	1070
1075	1	eu Thr Val Glu Glu 080	1085
1090	1095	la Gln Glu Glu Ile 110	0
	Glu Asp Leu G	lu Leu Ser Asn Lys 1115	Arg His Ser Leu 1120
1105 Val Gln Thr Leu		et Lys Arg Lys Leu 1130	
Ala Phe Val Gly	Gly Ser Arg A	la Ile Ile Leu Asp 1145	
	Tyr Ala Arg A	rg Ala Ile Trp Asp 160	
		eu Leu Ser Thr His 118	His Met Asp Glu
		le Ala Ile Ile Ser	His Gly Lys Leu
1185	1190	1195 he Leu Lys Gly Thr	1200
	1205	1210	1215
Tyr Arg Leu Thr 122		rg Pro Ala Glu Pro 1225	1230
	Ala Ser Ser P	ro Pro Gly Arg Ala 240	Pro Leu Ser Ser 1245
	Gln Val Ser G 1255	ln Phe Ile Arg Lys 126	
Cys Leu Leu Val		er Thr Glu Leu Ser	
1265	1270	1275	1280
	1285	la Phe Glu Arg Leu 1290	1295
130	0	is Leu Ser Ser Phe 1305	1310
1315	1	eu Lys Val Ser Glu 320	1325
1330	1335	al Lys Glu Ser Arg 134	0
1345	1350	er Gly Glu Gly His 1355	1360
Ala Arg Cys Ser	Glu Leu Thr G 1365	In Ser Gln Ala Ser 1370	Leu Gln Ser Ala 1375
Ser Ser Val Gly 138		ly Asp Glu Gly Ala 1385	Gly Tyr Thr Asp 1390
		eu Phe Asp Asn Pro	Gln Asp Pro Asp
	Gln Glu Val G	400 Iu Ala Glu Ala Leu	
1410	1415	142 ly Gly Trp Leu Lys	
1425	1430	1435	1440
His Gly Leu Leu	Val Lys Arg P 1445	he His Cys Ala Arg 1450	Arg Asn Ser Lys 1455
Ala Leu Phe Ser 146		eu Pro Ala Phe Phe 1465	Val Cys Val Ala 1470
Met Thr Val Ala 1475		ro Glu Ile Gly Ası 480	Leu Pro Pro Leu 1485
1490	1495	is Asn Tyr Thr Gli 150	0
		lu Arg Arg Glu Ty	
1505 Ser Bro Asp Ala	1510 Ser Pro Gln G	1515 In Leu Val Ser Thi	1520 The Arg Leu Pro
Ser Ero web wra	1525	1530	1535

```
Ser Gly Val Gly Ala Thr Cys Val Leu Lys Ser Pro Ala Asn Gly Ser
                                     1550
                   1545
        1540
Leu Gly Pro Thr Leu Asn Leu Ser Ser Gly Glu Ser Arg Leu Leu Ala
                                            1565
                       1560
Ala Arg Phe Phe Asp Ser Met Cys Leu Glu Ser Phe Thr Gln Gly Leu
                                         1580
                     1575
Pro Leu Ser Asn Phe Val Pro Pro Pro Pro Ser Pro Ala Pro Ser Asp
                                      1595
                  1590
Ser Pro Ala Ser Pro Asp Glu Asp Leu Gln Ala Trp Asn Val Ser Leu
                                  1610
              1605
Pro Pro Thr Ala Gly Pro Glu Met Trp Thr Ser Ala Pro Ser Leu Pro
                             1625
                                                 1630
           1620
Arg Leu Val Arg Glu Pro Val Arg Cys Thr Cys Ser Ala Gln Gly Thr
                                             1645
                          1640
      1635
Gly Phe Ser Cys Pro Ser Ser Val Gly Gly His Pro Pro Gln Met Arg
                      1655
                                         1660
Val Val Thr Gly Asp Ile Leu Thr Asp Ile Thr Gly His Asn Val Ser
                  1670
                                      1675
Glu Tyr Leu Leu Phe Thr Ser Asp Arg Phe Arg Leu His Arg Tyr Gly
                                  1690
                                                     1695
               1685
Ala Ile Thr Phe Gly Asn Val Leu Lys Ser Ile Pro Ala Ser Phe Gly
           1700
                              1705
                                                 1710
Thr Arg Ala Pro Pro Met Val Arg Lys Ile Ala Val Arg Arg Ala Ala
                          1720
                                             1725
       1715
Gln Val Phe Tyr Asn Asn Lys Gly Tyr His Ser Met Pro Thr Tyr Leu
                                         1740
                      1735
Asn Ser Leu Asn Asn Ala Ile Leu Arg Ala Asn Leu Pro Lys Ser Lys
                  1750
                                      1755
1745
Gly Asn Pro Ala Ala Tyr Gly Ile Thr Val Thr Asn His Pro Met Asn
                                  1770
               1765
Lys Thr Ser Ala Ser Leu Ser Leu Asp Tyr Leu Leu Gln Gly Thr Asp
                              1785
           1780
Val Val Ile Ala Ile Phe Ile Ile Val Ala Met Ser Phe Val Pro Ala
                          1800
       1795
Ser Phe Val Val Phe Leu Val Ala Glu Lys Ser Thr Lys Ala Lys His
                                         1820
                      1815
Leu Gln Phe Val Ser Gly Cys Asn Pro Ile Ile Tyr Trp Leu Ala Asn
                                     1835
                   1830
Tyr Val Trp Asp Met Leu Asn Tyr Leu Val Pro Ala Thr Cys Cys Val
                                 1850
              1845
Ile Ile Leu Phe Val Phe Asp Leu Pro Ala Tyr Thr Ser Pro Thr Asn
                                                1870
                             1865
           1860
Phe Pro Ala Val Leu Ser Leu Phe Leu Leu Tyr Gly Trp Ser Ile Thr
                                             1885
                         1880
       1875
Pro Ile Met Tyr Pro Ala Ser Phe Trp Phe Glu Val Pro Ser Ser Ala
                      1895
                                         1900
Tyr Val Phe Leu Ile Val Ile Asn Leu Phe Ile Gly Ile Thr Ala Thr
                                      1915
                  1910
Val Ala Thr Phe Leu Leu Gln Leu Phe Glu His Asp Lys Asp Leu Lys
               1925
                                  1930
Val Val Asn Ser Tyr Leu Lys Ser Cys Phe Leu Ile Phe Pro Asn Tyr
                              1945
                                                 1950
           1940
Asn Leu Gly His Gly Leu Met Glu Met Ala Tyr Asn Glu Tyr Ile Asn
                          1960
                                             1965
       1955
Glu Tyr Tyr Ala Lys Ile Gly Gln Phe Asp Lys Met Lys Ser Pro Phe
                      1975
                                         1980
   1970
Glu Trp Asp Ile Val Thr Arg Gly Leu Val Ala Met Ala Val Glu Gly
                                     1995
                   1990
Val Val Gly Phe Leu Leu Thr Ile Met Cys Gln Tyr Asn Phe Leu Arg
                           2010
              2005
Arg Pro Gln Arg Met Pro Val Ser Thr Lys Pro Val Glu Asp Asp Val
```

Asp Val Ala Ser Glu Arg Gln Arg Val Leu Arg Gly Asp Ala Asp Asn Asp Met Val Lys Ile Glu Asn Leu Thr Lys Val Tyr Lys Ser Arg Lys Ile Gly Arg Ile Leu Ala Val Asp Arg Leu Cys Leu Gly Val Arg Pro Gly Glu Cys Phe Gly Leu Leu Gly Val Asn Gly Ala Gly Lys Thr Ser Thr Phe Lys Met Leu Thr Gly Asp Glu Ser Thr Thr Gly Gly Glu Ala Phe Val Asn Gly His Ser Val Leu Lys Glu Leu Leu Gln Val Gln Gln Ser Leu Gly Tyr Cys Pro Gln Cys Asp Ala Leu Phe Asp Glu Leu Thr Ala Arg Glu His Leu Gln Leu Tyr Thr Arg Leu Arg Gly Ile Ser Trp Lys Asp Glu Ala Arg Val Val Lys Trp Ala Leu Glu Lys Leu Glu Leu Thr Lys Tyr Ala Asp Lys Pro Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr Ala Ile Ala Leu Ile Gly Tyr Pro Ala Phe Ile Phe Leu Asp Glu Pro Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Leu Ile Leu Asp Leu Ile Lys Thr Gly Arg Ser Val Val Leu Thr Ser His Ser Met Glu Glu Cys Glu Ala Leu Cys Thr Arg Leu Ala Ile Met Val Asn Gly Arg Leu Arg Cys Leu Gly Ser Ile Gln His Leu Lys Asn Arg Phe Gly Asp Gly Tyr Met Ile Thr Val Arg Thr Lys Ser Ser Gln Ser Val Lys Asp Val Val Arg Phe Phe Asn Arg Asn Phe Pro Glu Ala Met Leu Lys Glu Arg His His Thr Lys Val Gln Tyr Gln Leu Lys Ser Glu His Ile Ser Leu Ala Gln Val Phe Ser Lys Met Glu Gln Val Ser Gly Val Leu Gly Ile Glu Asp Tyr Ser Val Ser Gln Thr Thr Leu Asp Asn Val Phe Val Asn Phe Ala Lys Lys Gln Ser Asp Asn Leu Glu Gln Gln Glu Thr Glu Pro Pro Ser Ala Leu Gln Ser Pro Leu Gly Cys Leu Leu Ser Leu Leu Arg Pro Arg Ser Ala Pro Thr Glu Leu Arg Ala Leu Val Ala Asp Glu Pro Glu Asp Leu Asp Thr Glu Asp Glu Gly Leu Ile Ser Phe Glu Glu Glu Arg Ala Gln Leu Ser Phe Asn Thr Asp Thr Leu Cys

<210> 3

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223	3> PCR Primer	
<400)> 3	
	acgactcact atagggc	27
<210		
	L> 27 2> DNA	
	<pre>2> DNA 3> Artificial Sequence</pre>	
\21 3	37 Architetal Sequence	
<220) >	
<223	3> PCR Primer	
<400)> 4	
tgagtttgt	c cacgcagaca accagag	27
<210		
	0> 5 L> 23	
	2> DNA	
	3> Artificial Sequence	
\21.	> WICHITCHE Deficence	
<220)>	
	> PCR Primer	
<400)> 5	
actcactate	gggetegage gge	23
<210		
	> 23	
	?> DNA 3> Artificial Sequence	
\21.	> viciliorar podrence	
<220) >	
<223	3> PCR Primer	
<400	o> 6	
ccagctccad	tcccaggctt ctg	23
-01/)	
<210 <211	0> / L> 23	
	>> DNA	
	3> Artificial Sequence	
122		
<220) >	
<223	3> PCR Primer	
<400		~ ~
ccactgggc	a gcgagaagtt gtc	23
221)> 8	
	L> 23	
	2> DNA	
	3> Artificial Sequence	
	-	
<220		
<223	3> PCR Primer	

<4UU> 0		
gaagctggag ttctggcgga	tct	23
.010. 0		
<210> 9		
<211> 27		
<212> DNA		
<213> Artificial	Sequence	
	_	
<220>		
<223> PCR Primer		
<400> 9		
cagaccacac tggacaatgt	gttcgtg	27
<210> 10		
<211> 23		
<212> DNA		
<213> Artificial	sequence	
<220>		
<223> PCR Primer	•	
<400> 10		
tcatcagctt cgaggaggag	aaa	23
ccaccagect cyayyayyay	cgg	
<210> 11		
<211> 29		
<212> DNA		
<213> Artificial	Sequence	
<220>		
<223> PCR Primer	•	
<400> 11		
ataagcttgc tgaggcggcg	gagcgtggc	29
<210> 12		
<211> 20		
<212> DNA	_	•
<213> Artificial	Sequence	
<220>		
<223> PCR Primer	•	
<400> 12		
		20
ccactgggcg agaagttgtc		20
<210> 13		
<211> 20		
<212> DNA		
<213> Artificial	Semience	
(213) ALCILICIAI	. Pedremca	
<220>		
<2223 DCR Primer		

cctca	<400> 13 ttttc ccctacaacc	20
	<210> 14	
	<211> 26	
	<212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> PCR Primer	
	<400> 14	
acctg	ctcca tcttgctgct gaacac	26
	.010. 15	
	<210> 15 <211> 21	
	<211> 21 <212> DNA	
	<213> Artificial Sequence	
	12137 Militarian boquonoo	
	<220>	
	<223> PCR Primer	
	<400> 15	
cagcg	gcggc aacaagcgga a	21
	<210> 16	
	<211> 30	
	<212> DNA	
	<213> Artificial Sequence	
	<220>	
	<223> PCR Primer	
	<400> 16	
ggtga	attcg gcaggcactg ggggacttgt	30
	<210> 17	
	<211> 28	
	<211> 20 <212> DNA	
	<213> Artificial Sequence	
	1220 Indiana Doğumuy	
	<220>	
	<223> PCR Primer	
	<400> 17	
tagta	ctcct tgggcttcct gcaccagc	28
5		
	<210> 18	
	<211> 18 <211> 24	
	<211> 24 <212> DNA	
	<212> DNA <213> Artificial Sequence	
	THEOR INCHES DOGSTON	
	<220>	
	<223> PCR Primer	
	<400> 18	
	EAVU2 10	

ccagggcaga tgagggacca aaga	24
<210> 19	
<211> 23	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> PCR Primer	
<400> 19	
catccagcag gtcccccaga agc	23
<210> 20	
<211> 36	
<212> DNA	
<213> Artificial Sequence	
<220>	
<221> misc_feature	•
<222> (24, 25, 29, 30, 34, 35)	
<223> n = Inosine (I)	
(223) II = 11031110 (2)	
<400> 20	
ggccacgcgt cgactagtac gggnngggnn gggnng	36
ggccacgege cgaceageac gggggg	
<210> 21	
<211> 25	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> PCR Primer	
<400> 21	
aaacaggttg cccttcctcc accac	25
<210> 22	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> PCR Primer	
<400> 22	
ggccacgcgt cgactagtac	20
2200002020 020002000	
<210> 23	
<211> 22	
<211> 22 <212> DNA	
<212> DNA <213> Artificial Sequence	
(ST2) Wifilings befremes	
.000	
<220>	
ZZZKS DUK DTIMBT	

```
<400> 23
                                                                        22
acagcgattg catgacaggc ag
     <210> 24
     <211> 21
     <212> DNA
     <213> Artificial Sequence
     <220>
     <223> PCR Primer
     <400> 24
                                                                        21
agggagctgg ctacaccgac g
     <210> 25
     <211> 21
     <212> DNA
     <213> Artificial Sequence
     <220>
     <223> PCR Primer
     <400> 25
                                                                        21
cgcctgtgac cacccgcatc t
     <210> 26
     <211> 188
     <212> PRT
     <213> Homo Sapiens
     <400> 26
Ala Val Asp Gly Leu Ala Leu Asn Phe Tyr Glu Gly Gln Ile Thr Ser
                 5
                                    10
1
Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu
                                25
Thr Gly Leu Phe Pro Pro Thr Ser Gly Thr Ala Tyr Ile Leu Gly Lys
                           40
Asp Ile Arg Ser Glu Met Ser Thr Ile Arg Gln Asn Leu Gly Val Cys
                        55
Pro Gln His Asn Val Leu Phe Asp Met Leu Thr Val Glu Glu His Ile
                                        75
                    70
Trp Phe Tyr Ala Arg Leu Lys Gly Leu Ser Glu Lys His Val Lys Ala
                                    90
               85
Glu Met Glu Gln Met Ala Leu Asp Val Gly Leu Pro Ser Ser Lys Leu
                                105
Lys Ser Lys Thr Ser Gln Leu Ser Gly Gly Met Gln Arg Lys Leu Ser
                            120
Val Ala Leu Ala Phe Val Gly Gly Ser Lys Val Val Ile Leu Asp Glu
                                            140
                        135
Pro Thr Ala Gly Val Asp Pro Tyr Ser Arg Arg Gly Ile Trp Glu Leu
                                        155
                    150
Leu Leu Lys Tyr Arg Gln Gly Arg Thr Ile Ile Leu Ser Thr His His
                                    170
Met Asp Glu Ala Asp Val Leu Gly Asp Arg Ile Ala
```

<210> 27

185

<211> 183 <212> PRT

<213> Homo Sapiens

<400> 27 Ala Val Asp Arg Ile Cys Val Gly Ile Pro Pro Gly Glu Cys Phe Gly 10 Leu Leu Gly Val Asn Gly Ala Gly Lys Ser Ser Thr Phe Lys Met Leu 25 20 Thr Gly Asp Thr Thr Val Thr Arg Gly Asp Ala Phe Leu Asn Arg Asn 40 Ser Ile Leu Ser Asn Ile His Glu Val His Gln Asn Met Gly Tyr Cys 55 Pro Gln Phe Asp Ala Ile Thr Glu Leu Leu Thr Gly Arg Glu His Val 75 70 Glu Phe Phe Ala Leu Leu Arg Gly Val Pro Glu Lys Glu Val Gly Lys 90 85 Val Gly Glu Trp Ala Ile Arg Lys Leu Gly Leu Val Lys Tyr Gly Glu 105 110 100 Lys Tyr Ala Gly Asn Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr 120 115 Ala Met Ala Leu Ile Gly Gly Pro Pro Val Val Phe Leu Asp Glu Pro 135 Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Cys Ala 150 155 Leu Ser Val Val Lys Glu Gly Arg Ser Val Val Leu Thr Ser His Ser 170 165 Met Glu Glu Cys Glu Ala Leu 180

<210> 28

<211> 187

<212> PRT

<213> Mus musculus

180

<400> 28 Ala Leu Asn Lys Leu Ser Leu Asn Leu Tyr Glu Asn Gln Val Val Ser Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu 25 Thr Gly Leu Phe Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His 40 45 Asp Ile Arg Thr Glu Met Asp Glu Ile Arg Lys Asn Leu Gly Met Cys 55 60 Pro Gln His Asn Val Leu Phe Asp Arg Leu Thr Val Glu Glu His Leu 70 75 Trp Phe Tyr Ser Arg Leu Lys Ser Met Ala Gln Glu Glu Ile Arg Lys 90 85 Glu Thr Asp Lys Met Ile Glu Asp Leu Glu Leu Ser Asn Lys Arg His 105 100 Ser Leu Val Gln Thr Leu Ser Gly Gly Met Lys Arg Lys Leu Ser Val 120 115 Ala Ile Ala Phe Val Gly Gly Ser Arg Ala Ile Ile Leu Asp Glu Pro 135 140 Thr Ala Gly Val Asp Pro Tyr Ala Arg Arg Ala Ile Trp Asp Leu Ile 155 150 Leu Lys Tyr Lys Pro Gly Arg Thr Ile Leu Leu Ser Thr His His Met 170 165 Asp Glu Ala Asp Leu Leu Gly Asp Arg Ile Ala

185

<210> 29 <211> 184 <212> PRT <213> Mus musculus

<400> 29 Ala Val Asp Arg Leu Cys Leu Gly Val Cys Val Pro Gly Glu Cys Phe 5 10 Gly Leu Leu Gly Val Asn Gly Ala Gly Lys Thr Ser Thr Phe Lys Met 25 Leu Thr Gly Asp Glu Ser Thr Thr Gly Gly Glu Ala Phe Val Asn Gly 40 His Ser Val Leu Lys Asp Leu Leu Gln Val Gln Gln Ser Leu Gly Tyr 55 Cys Pro Gln Phe Asp Ala Leu Phe Asp Glu Leu Thr Ala Arg Glu His 75 70 Leu Gln Leu Tyr Thr Arg Leu Arg Gly Ile Pro Trp Lys Asp Glu Ala 90 85 Gln Val Val Lys Trp Ala Leu Glu Lys Leu Glu Leu Thr Lys Tyr Ala 105 100 Asp Lys Pro Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser 120 115 Thr Ala Ile Ala Leu Ile Gly Tyr Pro Ala Phe Ile Phe Leu Asp Glu 135 Pro Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Leu 155 150 Ile Leu Asp Leu Ile Lys Thr Gly Arg Ser Val Val Leu Thr Ser His 165 170 Ser Met Glu Glu Cys Glu Ala Leu 180

<210> 30 <211> 187 <212> PRT <213> Homo sapiens

<400> 30 Ala Leu Asn Lys Leu Ser Leu Asn Leu Tyr Glu Asn Gln Val Val Ser 10 5 1 Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu 25 20 Thr Gly Leu Phe Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His 40 35 Asp Ile Arg Thr Glu Met Asp Glu Ile Arg Lys Asn Leu Gly Met Cys 55 60 Pro Gln His Asn Val Leu Phe Asp Arg Leu Thr Val Glu Glu His Leu 70 75 Trp Phe Tyr Ser Arg Leu Lys Ser Met Ala Gln Glu Glu Ile Arg Arg 90 Glu Met Asp Lys Met Ile Glu Asp Leu Glu Leu Ser Asn Lys Arg His 105 Ser Leu Val Gln Thr Leu Ser Gly Gly Met Lys Arg Lys Leu Ser Val 125 120 Ala Ile Ala Phe Val Gly Gly Ser Arg Ala Ile Ile Leu Asp Glu Pro 135 140 Thr Ala Gly Val Asp Pro Tyr Ala Arg Arg Ala Ile Trp Asp Leu Ile 155 150 Leu Lys Tyr Lys Pro Gly Arg Thr Ile Leu Leu Ser Thr His His Met 165 170 175
Asp Glu Ala Asp Leu Leu Gly Asp Arg Ile Ala
180 185

<210> 31 <211> 183 <212> PRT <213> Homo sapiens

<400> 31 Ala Val Asp Arg Leu Cys Leu Gly Val Arg Pro Gly Glu Cys Phe Gly 10 Leu Leu Gly Val Asn Gly Ala Gly Lys Thr Ser Thr Phe Lys Met Leu 20 25 Thr Gly Asp Glu Ser Thr Thr Gly Glu Ala Phe Val Asn Gly His 35 40 Ser Val Leu Lys Glu Leu Leu Gln Val Gln Gln Ser Leu Gly Tyr Cys 55 Pro Gln Cys Asp Ala Leu Phe Asp Glu Leu Thr Ala Arg Glu His Leu 75 Gln Leu Tyr Thr Arg Leu Arg Gly Ile Ser Trp Lys Asp Glu Ala Arg 85 90 Val Val Lys Trp Ala Leu Glu Lys Leu Glu Leu Thr Lys Tyr Ala Asp 100 105 110 Lys Pro Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr 120 125 Ala Ile Ala Leu Ile Gly Tyr Pro Ala Phe Ile Phe Leu Asp Glu Pro 135 140 Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn Leu Ile 155 150 Leu Asp Leu Ile Lys Thr Gly Arg Ser Val Val Leu Thr Ser His Ser 170 165 Met Glu Glu Cys Glu Ala Leu

<210> 32 <211> 187 <212> PRT <213> Homo sapiens

180

<400> 32 Ala Val Arg Asp Leu Asn Leu Asn Leu Tyr Glu Gly Gln Ile Thr Val 5 10 Leu Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Leu Ser Met Leu 25 Thr Gly Leu Phe Pro Pro Thr Ser Gly Arg Ala Tyr Ile Ser Gly Tyr 40 45 Glu Ile Ser Gln Asp Met Val Gln Ile Arg Lys Ser Leu Gly Leu Cys 55 60 Pro Gln His Asp Ile Leu Phe Asp Asn Leu Thr Val Ala Glu His Leu 70 75 Tyr Phe Tyr Ala Gln Leu Lys Gly Leu Ser Arg Gln Lys Cys Pro Glu 90 85 Glu Val Lys Gln Met Leu His Ile Ile Gly Leu Glu Asp Lys Trp Asn 105 Ser Arg Ser Arg Phe Leu Ser Gly Gly Met Arg Arg Lys Leu Ser Ile 120 Gly Ile Ala Leu Ile Ala Gly Ser Lys Val Leu Ile Leu Asp Glu Pro 130 135

<210> 33 <211> 183 <212> PRT <213> Homo sapiens

<400> 33 Ala Val Asp Arg Leu Ser Leu Ala Val Gln Lys Gly Glu Cys Phe Gly 10 5 Leu Leu Gly Phe Asn Gly Ala Gly Lys Thr Thr Thr Phe Lys Met Leu 25 20 Thr Gly Glu Glu Ser Leu Thr Ser Gly Asp Ala Phe Val Gly Gly His 40 Arg Ile Ser Ser Asp Val Gly Lys Val Arg Gln Arg Ile Gly Tyr Cys 55 Pro Gln Phe Asp Ala Leu Leu Asp His Met Thr Gly Arg Glu Met Leu 70 Val Met Tyr Ala Arg Leu Arg Gly Ile Pro Glu Arg His Ile Gly Ala 90 95 Cys Val Glu Asn Thr Leu Arg Gly Leu Leu Glu Pro His Ala Asn 110 105 100 Lys Leu Val Arg Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr 125 115 120 Gly Ile Ala Leu Ile Gly Glu Pro Ala Val Ile Phe Leu Asp Glu Pro 140 135 Ser Thr Gly Met Asp Pro Val Ala Arg Arg Leu Leu Trp Asp Thr Val 150 155 Ala Arg Ala Arg Glu Ser Gly Lys Ala Ile Ile Ile Thr Ser His Ser 170 165 Met Glu Glu Cys Glu Ala Leu 180

<210> 34 <211> 187 <212> PRT <213> Homo sapiens

<400> 34 Ala Val Asp Arg Leu Asn Ile Thr Phe Tyr Glu Asn Gln Ile Thr Ala 1 10 Phe Leu Gly His Asn Gly Ala Gly Lys Thr Thr Thr Leu Ser Ile Leu 25 30 Thr Gly Leu Leu Pro Pro Thr Ser Gly Thr Val Leu Val Gly Gly Arg 40 45 Asp Ile Glu Thr Ser Leu Asp Ala Val Arg Gln Ser Leu Gly Met Cys 60 55 Pro Gln His Asn Ile Leu Phe His His Leu Thr Val Ala Glu His Met 75 70 Leu Phe Tyr Ala Gln Leu Lys Gly Lys Ser Gln Glu Glu Ala Gln Leu 90 85 Glu Met Glu Ala Met Leu Glu Asp Thr Gly Leu His His Lys Arg Asn 100 105 Glu Glu Ala Gln Asp Leu Ser Gly Gly Met Gln Arg Lys Leu Ser Val Ala Ile Ala Phe Val Gly Asp Ala Lys Val Val Ile Leu Asp Glu Pro
130

Thr Ser Gly Val Asp Pro Tyr Ser Arg Arg Ser Ile Trp Asp Leu Leu
145

Leu Lys Tyr Arg Ser Gly Arg Thr Ile Ile Met Ser Thr His His Met
165

Asp Glu Ala Asp Leu Leu Gly Asp Arg Ile Ala
180

<210> 35 <211> 183 <212> PRT

<213> Homo Sapiens

<400> 35 Ala Val Asp Arg Leu Cys Val Gly Val Arg Pro Gly Glu Cys Phe Gly 10 Leu Leu Gly Val Asn Gly Ala Gly Lys Thr Thr Thr Phe Lys Met Leu 25 20 Thr Gly Asp Thr Thr Val Thr Ser Gly Asp Ala Thr Val Ala Gly Lys 40 35 Ser Ile Leu Thr Asn Ile Ser Glu Val His Gln Asn Met Gly Tyr Cys 55 Pro Gln Phe Asp Ala Ile Asp Glu Leu Leu Thr Gly Arg Glu His Leu Tyr Leu Tyr Ala Arg Leu Arg Gly Val Pro Ala Glu Glu Ile Glu Lys 90 Val Ala Asn Trp Ser Ile Lys Ser Leu Gly Leu Thr Val Tyr Ala Asp 110 105 Cys Leu Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu Ser Thr 125 120 Ala Ile Ala Leu Ile Gly Cys Pro Pro Leu Val Leu Leu Asp Glu Pro 140 135 Thr Thr Gly Met Asp Pro Gln Ala Arg Arg Met Leu Trp Asn Val Ile 155 150 Val Ser Ile Ile Arg Lys Gly Arg Ala Val Val Leu Thr Ser His Ser 170 165 Met Glu Glu Cys Glu Ala Leu 180

<210> 36 <211> 13 <212> PRT <213> Homo sapiens

<400> 36
Gly Gln Ser Arg Lys Leu Asp Gly Gly Trp Leu Lys Val
1 5 10